

Figure legends:

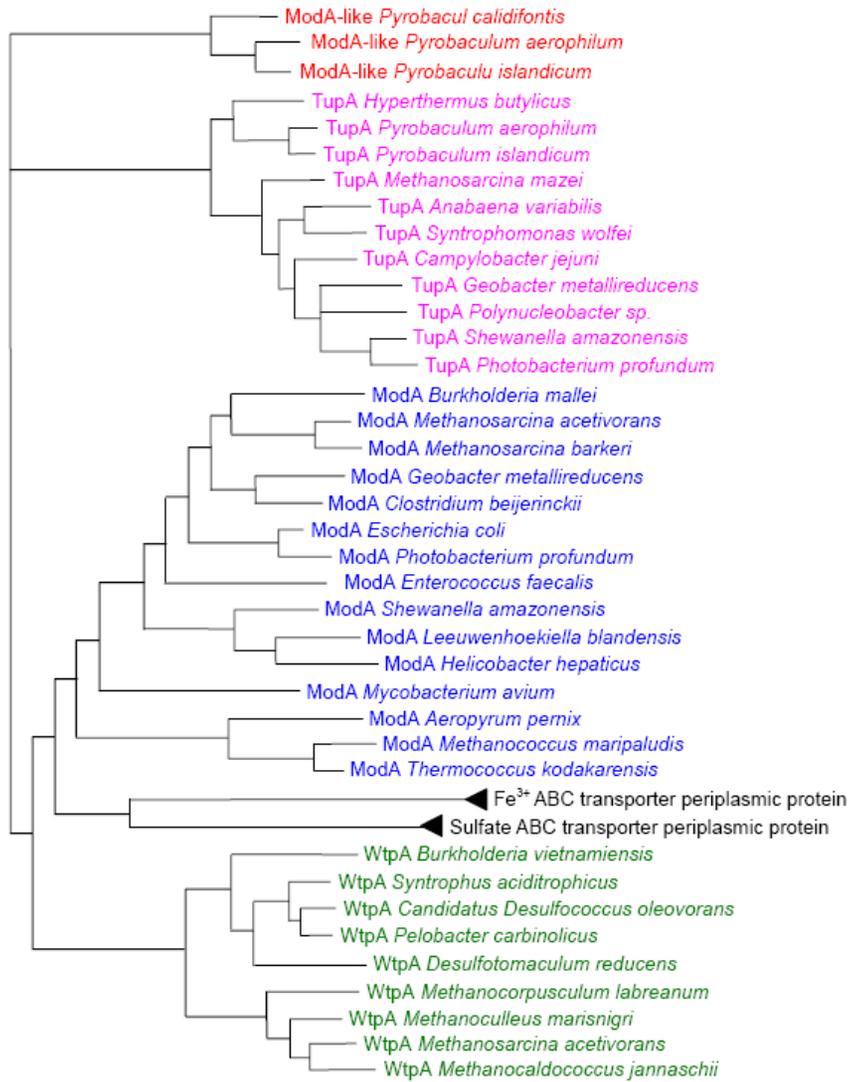
Fig. S1. Phylogenetic trees of periplasmic components of prokaryotic Mo/W transporters based on MrBayes and PHYML. ModA-like proteins are shown in red and bold, TupA in pink, WtpA in green and ModA in blue. The sulfate and Fe³⁺ ABC transporter branches were compressed and represented by family names. These trees were generated by corresponding programs and visualized via TreeView program. A. MrBayes analysis; B. PHYML analysis.

Fig. S2. CLUSTALW multiple alignment of ModA, WtpA and ModA-like. Residues involved in molybdate binding in ModA are shown in red background and residues involved in tungstate binding in WtpA in blue background. Overlapped residues between ModA and WtpA are shown in red background. Other residues shown in white on black or grey are conserved in homologues. The secondary structures of ModA and WtpA are based on the crystal structures identified in *E. coli* and *Archaeoglobus fulgidus*, respectively. Secondary structures of ModA-like proteins were predicted using GOR IV with default parameters (http://npsa-pbil.ibcp.fr/cgi-bin/npsa_automat.pl?page=npsa_gor4.html).

Fig. S3. MUSCLE multiple alignment of ModA, WtpA and ModA-like. See Fig. S2 legend for annotation of functional residues.

Fig. S1

A.



B.

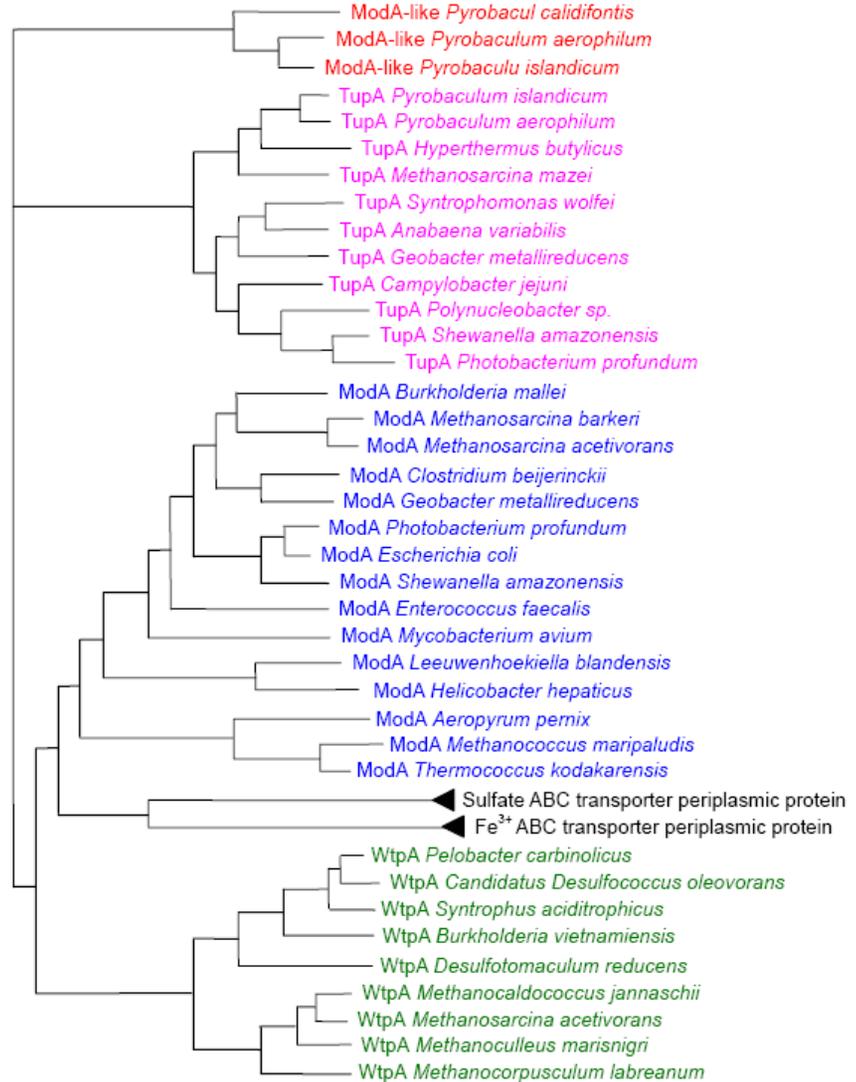
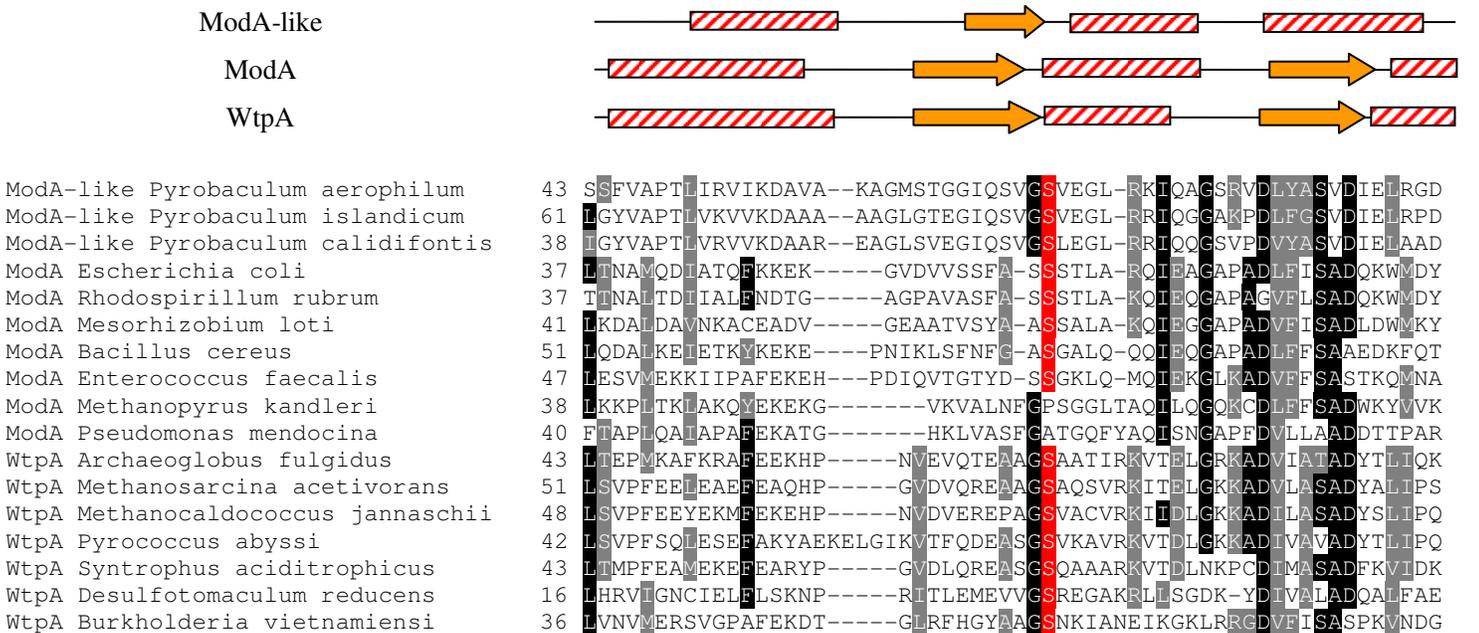
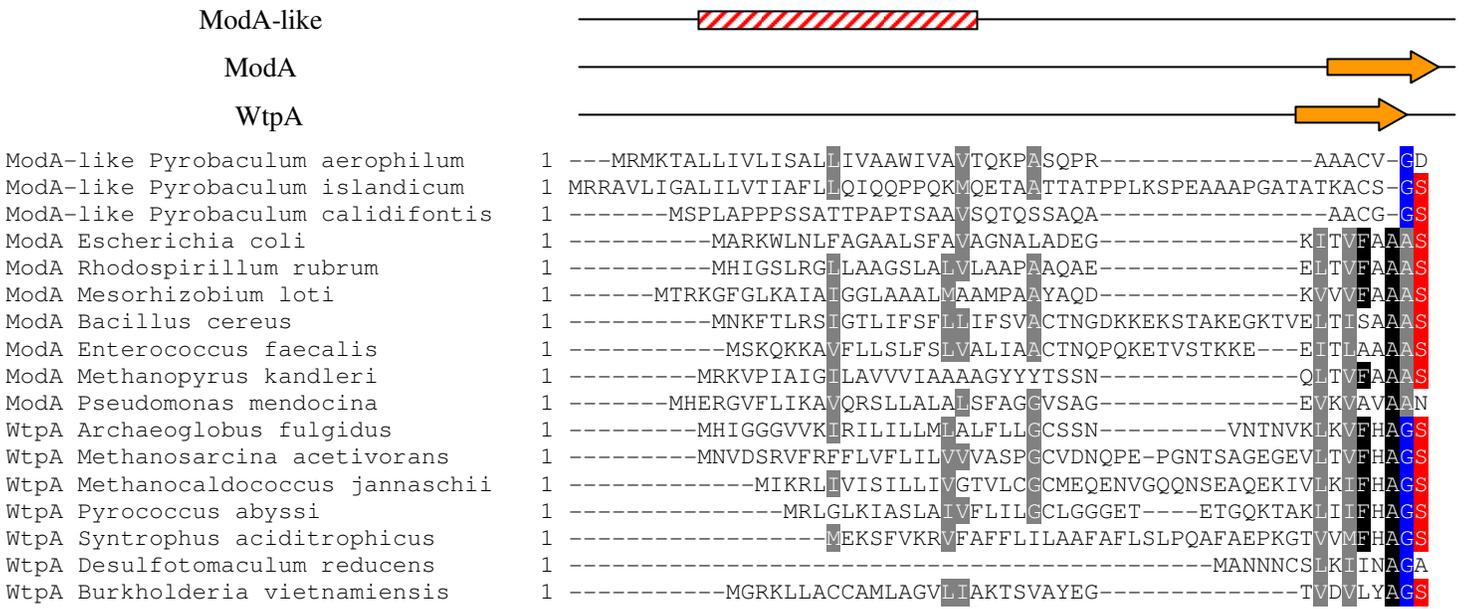


Fig. S2



ModA <i>Enterococcus faecalis</i>	212	I	V	A	A	M	P	---	E	A	V	L	K	K	P	I	I	M	P	V	G	K	V	A	A	S	K	K	Q	S	A	D	A	F	L	N	F	L	S	Q	Q	C	R	K	Y	F	E	N	I	G	F	K	L	T	K	---					
ModA <i>Methanopyrus kandleri</i>	202	I	V	Q	M	F	P	---	H	S	L	T	G	P	I	I	I	W	G	A	A	V	I	K	G	N	E	-	Q	L	A	E	D	F	L	N	Y	C	L	E	H	I	D	E	F	K	K	Y	G	W	S	P	A	-----							
ModA <i>Pseudomonas mendocina</i>	204	S	G	S	A	W	I	V	P	--	E	A	M	Y	E	P	I	R	Q	D	A	V	I	L	R	Q	G	A	G	N	P	A	A	A	A	L	V	E	Y	L	Q	G	A	E	A	A	R	V	I	E	S	F	G	Y	K	L	P	---			
WtpA <i>Archaeoglobus fulgidus</i>	272	L	A	N	G	---	K	E	V	T	G	K	P	I	V	Y	G	T	I	P	K	N	A	E	N	R	E	L	A	V	E	F	V	K	L	V	I	S	E	E	Q	E	I	L	R	-	E	L	G	Q	E	P	L	V	P	---					
WtpA <i>Methanosarcina acetivorans</i>	281	M	V	N	G	---	E	V	V	T	G	S	P	I	V	Y	G	V	T	I	P	N	N	A	E	N	S	E	L	A	T	E	F	V	A	L	L	G	E	T	G	Q	Q	I	F	L	-	E	N	G	Q	P	P	I	V	P	---				
WtpA <i>Methanocaldococcus jannaschii</i>	275	I	I	A	K	N	---	K	T	I	N	A	K	P	I	V	Y	G	M	T	V	P	T	N	A	P	H	K	E	A	I	E	F	V	K	F	V	L	G	--	H	P	E	V	L	E	-	N	N	G	Q	P	A	I	I	P	---				
WtpA <i>Pyrococcus abyssi</i>	271	L	G	S	T	---	K	T	I	Y	A	K	P	I	V	Y	G	T	I	V	L	K	D	A	P	N	R	E	L	A	L	E	F	L	K	F	L	S	E	K	G	K	E	I	F	R	-	E	N	H	Q	D	F	L	T	P	---				
WtpA <i>Syntrophus aciditrophicus</i>	246	V	T	G	K	K	P	G	E	Y	M	E	M	K	G	A	S	T	T	Y	G	I	T	L	I	K	N	A	P	N	Q	E	A	A	V	A	F	L	E	Y	M	L	N	P	K	G	G	L	K	I	L	K	D	Q	G	P	P	F	I	P	---
WtpA <i>Desulfotomaculum reducens</i>	218	V	E	S	K	I	P	G	K	N	I	I	H	G	K	P	T	E	E	A	V	G	L	V	K	T	T	Q	N	P	E	L	A	Q	A	F	I	D	L	L	A	G	P	E	G	-	H	A	I	L	E	E	C	G	---	L	I	P	---		
WtpA <i>Burkholderia vietnamiensis</i>	215	I	P	A	V	R	P	---	A	P	E	L	Q	A	K	A	R	Y	T	L	I	L	G	D	A	P	N	P	A	G	A	A	R	F	V	D	E	L	L	S	A	K	G	R	A	-	L	K	E	H	G	V	D	V	V	K	P	T	---		

ModA-like



ModA

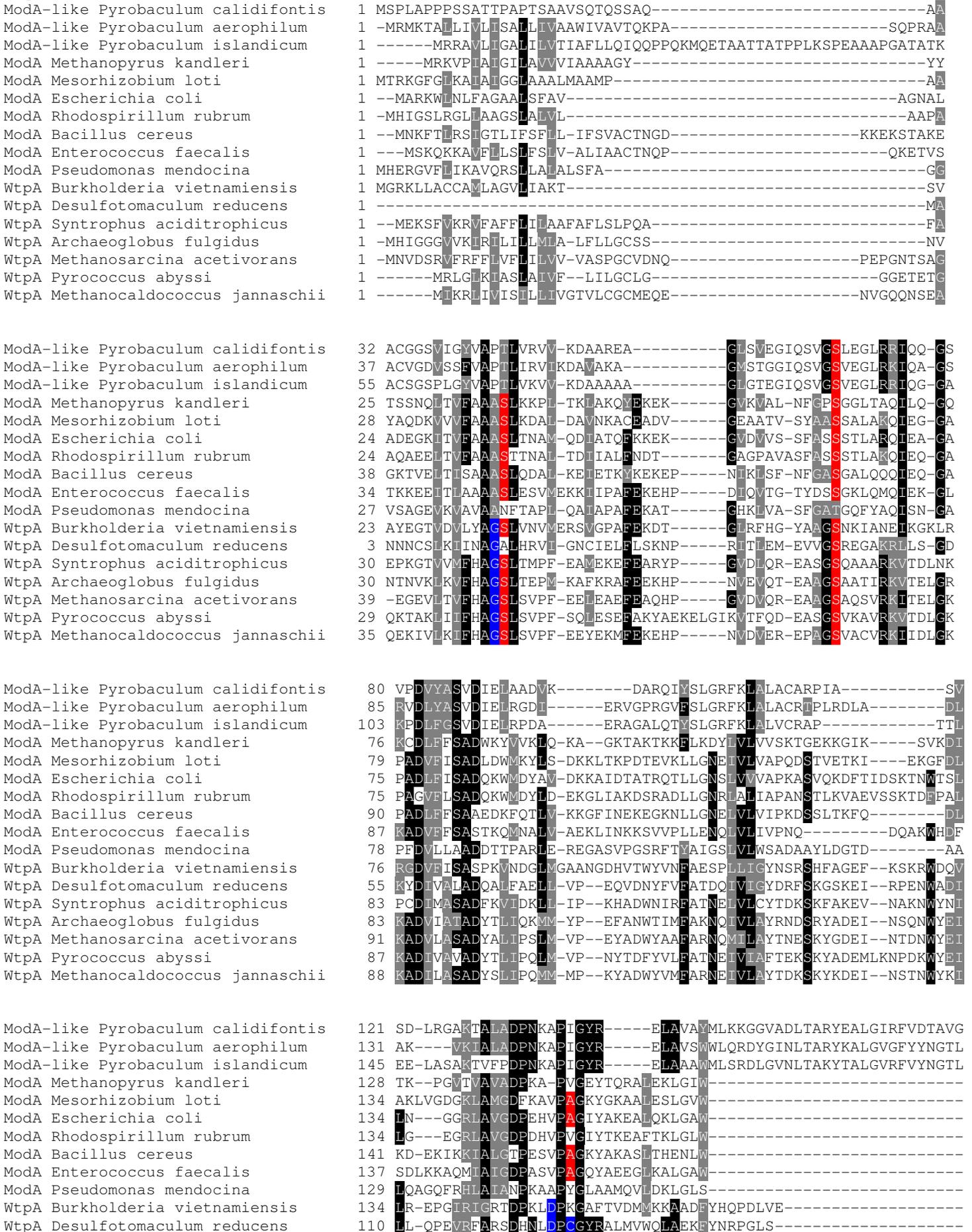


WtpA



ModA-like <i>Pyrobaculum aerophilum</i>	312	N	M	N	L	T	A	Y	G	F	V	K	-----													
ModA-like <i>Pyrobaculum islandicum</i>	325	K	M	D	L	S	K	Y	G	F	I	-----														
ModA-like <i>Pyrobaculum calidifontis</i>	305	K	M	D	L	A	K	Y	G	F	V	K	-----													
ModA <i>Escherichia coli</i>		-----																								
ModA <i>Rhodospirillum rubrum</i>		-----																								
ModA <i>Mesorhizobium loti</i>	264	N	-----																							
ModA <i>Bacillus cereus</i>		-----																								
ModA <i>Enterococcus faecalis</i>		-----																								
ModA <i>Methanopyrus kandleri</i>		-----																								
ModA <i>Pseudomonas mendocina</i>		-----																								
WtpA <i>Archaeoglobus fulgidus</i>	326	P	R	A	D	T	A	V	P	S	L	K	A	M	V	E	V	S	-----							
WtpA <i>Methanosarcina acetivorans</i>	335	A	I	A	E	G	K	D	S	M	P	E	E	L	Q	A	L	V	V	-----						
WtpA <i>Methanocaldococcus jannaschii</i>	328	A	V	A	Y	G	N	V	P	E	E	L	K	D	L	V	K	I	E	K	-----					
WtpA <i>Pyrococcus abyssi</i>	326	P	V	A	F	G	N	V	P	E	E	I	K	G	L	V	E	I	K	E	-----					
WtpA <i>Syntrophus aciditrophicus</i>	306	C	R	V	P	T	Q	K	M	K	S	T	L	P	E	K	L	Q	K	L	V	E	V	K	D	---
WtpA <i>Desulfotomaculum reducens</i>	274	C	-----																							
WtpA <i>Burkholderia vietnamiensis</i>	272	V	A	G	N	A	Q	A	V	P	P	S	L	Q	A	V	I	D	A	A	Q	-----				

Fig. S3



WtpA Syntrophus aciditrophicus 138 LQ-RKGVVWGHSDPNLDPCCGYRALMVLQLAEKYKPKGLY-----
WtpA Archaeoglobus fulgidus 138 LK-RPDVRFGFSNPNDPCGYRSLMATQLAEIYYNDPTIFDELVAKNSLRF-SEDNGSY
WtpA Methanosarcina acetivorans 146 LR-RPDVRYGFSNPNDPCAGYRSQMVTLQAESYYNDDMIYDDLMLANTGMTLTTEENGTA
WtpA Pyrococcus abyssi 144 LA-REDVSEGFSDPNQDPCGYRSVMVMKLDLYYGKP-IFETLVEKTTNIYA----NGTH
WtpA Methanocaldococcus jannaschii 143 LQ-RPDVKIGFSDNPNDPCGYRTQMVLQLAEIYYKDPTIYDNLVLKHSNIKV-EENNGTY

ModA-like Pyrobaculum calidifontis 175 FNVTV--PTALFSTAEVVKVAPSLDGSWSMFEITGOVDCVFAFLP-FILGKGVWELKP--AGA
ModA-like Pyrobaculum aerophilum 182 TITV---PSALPNTNITDLPASLDGAWTKLEIGAVDCVYAYVP-FLVNHLSLKP--HGE
ModA-like Pyrobaculum islandicum 199 YVYI---PSSLPNTEADVAPNLDGSAWKEEAGAGRCMFAYVP-FLLGKGLQLRP--AGG
ModA Methanopyrus kandleri 158 -----DKIVKNGNLKARPGTVNQVATMVKNDQIDAGFVYRS-VAVGFGLPIVQ--MFP
ModA Mesorhizobium loti 167 -----SSVEGK---VAQAEVRAALKLVSTGEAALGIYVATDAHADKGVKVVG--TFP
ModA Escherichia coli 164 -----DTLSPK---LAPAEVVRGALALVERNEAPLGIYVYGSDAVASKGVKVA--TFP
ModA Rhodospirillum rubrum 164 -----DGVANK---LAPANDVRGALTFVERGESPLGVVYSTDSAVSEKVKTLA--LPL
ModA Bacillus cereus 173 -----NDVQNK---FVFTKDVQRVLTIVYETIGNVDAGIVYKTDALISEKVKIGE--TAA
ModA Enterococcus faecalis 170 -----SYVEKH---ASFGTNVTEVLEWVANASAEAGLIVYATDAATNSKVAIYA--AMP
ModA Pseudomonas mendocina 162 -----EAVKKG---LVEGQNIQAQHFVSTGNAELGFVALSQVYQDGLGSGSAWIVP
WtpA Burkholderia vietnamiensis 174 -----KTLGAPENPEQVLPET-LVGRLOSQQLDAGFFVYST-ETSDLKIPAVR--PAP
WtpA Desulfotomaculum reducens 149 -----NELKACIPYTYPKSMDLAGALLEGKVDYAFIYSS-EAKQLGFPPYIN--LPS
WtpA Syntrophus aciditrophicus 177 -----EKLANRPVENRPKSVELVSLLOQTGNMDYAWVEYRS-VAVQENLKYVS--LPD
WtpA Archaeoglobus fulgidus 196 VLRMPSSERTEINKSKIMRSMEMELIHLVESGELDYFIYKS-VAKQHGFNVE--LPV
WtpA Methanosarcina acetivorans 205 LIHVPASEETSPNTSKIMRSMFVELSSALETGEIDYLYYRS-VAEQHGFEYVA--LPP
WtpA Pyrococcus abyssi 198 -IYAP--KELIVKDKRVVVRPKETDLVGLVESGLDYFIYKS-VAEQENLKYIT--LPN
WtpA Methanocaldococcus jannaschii 201 LILVP--KELDVDTNKLFVRSKETDLLAPLEAGAFDYFIYKS-VANQENLKYIE--LPK

ModA-like Pyrobaculum calidifontis 230 SAELWEVYVGDY-GRGQFYVYVFKPPYDFQDPLAIYVNFVDPSSGRVRAVVRVGRFEAF
ModA-like Pyrobaculum aerophilum 236 ETGYWTAYIGEAPGGKAVYVYVFKPPYDFLNDPPMPVRVLLDPSGKPAKINVGHFEEAF
ModA-like Pyrobaculum islandicum 253 GTSYWEPTYAEA-GGRYYVYVFKPPYDFAEDPPIKAVALLGP---PVKTIIRVGHFEEAF
ModA Methanopyrus kandleri 208 H-----SLTGPIIWAAGAVIK-GGNEQL
ModA Mesorhizobium loti 215 E-----DSHPPIIYPVAQTADSKDKD-
ModA Escherichia coli 212 E-----DSHKKVEYPAVVE-GHNNAT
ModA Rhodospirillum rubrum 212 E-----SSHKPVTYPVATVK-AKDTKT
ModA Bacillus cereus 221 A-----TSHEPIHYPLGVIKESKHKKE
ModA Enterococcus faecalis 218 E-----AVLKKPIIYPVGVKVAASKKQKS
ModA Pseudomonas mendocina 212 E-----AMYEPVIRQDAVILRQAGNPA
WtpA Burkholderia vietnamiensis 223 E-----LQAKARYTITLGDAPNPAG
WtpA Desulfotomaculum reducens 199 KINL-----SNPAHADYDQASISVESKIPGKNIITHGKPIEEAVGLVKTTONPEL
WtpA Syntrophus aciditrophicus 227 KINL-----GNYKEDPYYSQAVVKVTGKKPGEYMEKGASTTYGTLIKNAPNQEAF
WtpA Archaeoglobus fulgidus 253 EIDL-----SSPDYAEIYSKVKVVLANKG----EVTGKPIVYGITIPKNAENREL
WtpA Methanosarcina acetivorans 262 AIDL-----SSLEYADNYSKVQVEMVNGE----VVTGSPVIVYGVITPNNANSENEL
WtpA Pyrococcus abyssi 252 EINL-----KDFSKADFYKKSITLGTSGK----TYAKPIVYGVITVLKADAPNREL
WtpA Methanocaldococcus jannaschii 256 EINL-----GYEYADTYKKVALKIIAKNK----TINAKPIVYGVITVPTNAPHKKE

ModA-like Pyrobaculum calidifontis 289 VASFTERGDCVVEALKK-MDLAKYGFVK-----
ModA-like Pyrobaculum aerophilum 296 VASFTEKGDVLEALKN-MNLTAYGFVK-----
ModA-like Pyrobaculum islandicum 309 VASYTELGNVIESLKK-MDLASKYGFII-----
ModA Methanopyrus kandleri 229 AEDFL---NVCLEHIDE---FKKYGWSPA-----
ModA Mesorhizobium loti 236 TTAFL---KCLQSAKAG-ALFKEQGFVLTPSN-----
ModA Escherichia coli 233 VKAFY---DYLKGPQAA-EIFKRYGFTIK-----
ModA Rhodospirillum rubrum 233 ARAFV---DELKGAkak-AVFERYGFQVKP-----
ModA Bacillus cereus 243 ATSFY---EYLQSKDAQ-SLFFKRYGFTVLS-----
ModA Enterococcus faecalis 241 ADAFL---NFLQSQQCR-KYFENIGFKLTK-----
ModA Pseudomonas mendocina 234 AAALV---EYLGAEAA-RVIESFGYKLP-----
WtpA Burkholderia vietnamiensis 244 AARFV---DELLSAKGR-ALLKEHGVVVVKP---TVAGNAQAVPPSLQAVIDAAQ
WtpA Desulfotomaculum reducens 250 AQAFI---DILLAGPEGH-AILEECGLIPC-----
WtpA Syntrophus aciditrophicus 278 AVAFV---EYMLNPKGGLKILKDGQPPFIPCRVPTQKMKSTLPEKLQKLVVEVKD
WtpA Archaeoglobus fulgidus 299 AVEFV---KLVISEEQ-EILRELQGEPLVP---PRA---DTAVPSLKAMVEVS-
WtpA Methanosarcina acetivorans 308 ATEFV---ALLGETGQ-QIFIENGQPIVP---AIAEGKDSMPEELQALVV---
WtpA Pyrococcus abyssi 299 ALEFL---KFLISEKKG-EIFRENHQDFLTP---PVA---FGNVPEEIKGLVEIKE
WtpA Methanocaldococcus jannaschii 303 AIEFV---KEVILGH--P-EVLENNQPAIIP---AVA---YGNVPEELKDLVKIEK